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Hormonal regulation of leptin and leptin receptor expression in porcine subcutaneous adipose tissue^{1,2}

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ABSTRACT: The current study was performed to examine the response of the leptin gene to hormonal stimuli in porcine adipose tissue from finishing pigs. Yorkshire gilts (approximately 150 kg BW) were used in this study. Tissue from four to six pigs was used per experiment. Dorsal subcutaneous adipose tissue samples were acquired, and adipose tissue explants (approximately 100 mg) were prepared using sterile technique. Tissue slices were transferred to 12-well tissue culture plates containing 1 mL of Media 199 with 25 mM HEPES, 0.5% BSA, pH 7.4, and various hormone supplements. Triplicate tissue slices were incubated with either basal medium or hormone-supplemented media in a tissue culture incubator at 37°C with 95% air:5% CO₂. Hormones included insulin (100 nM), dexamethasone (1 μM), porcine GH, 100 ng/mL, triiodothyronine (T₃, 10 nM), porcine leptin (100 ng/mL), or IGF-I (250 ng/mL). Following incubation for 24 h, tissue samples from the incubations were blotted and transferred to microfuge tubes, frozen in liquid N, and stored at -80°C before analysis for gene mRNA abundance by reverse-transcription PCR and subsequent quantification of transcripts by capillary electrophoresis with laser-induced fluorescence detection. Media from the in-

cubations were collected in microfuge vials and stored at -20°C before analysis for leptin content by RIA. Insulin was required to maintain tissue and mRNA integrity; therefore, insulin was included in all incubations. The combination of insulin and dexamethasone stimulated leptin secretion into the medium by 60% ($P < 0.05$; $n = 6$). Porcine GH inhibited insulin induced leptin secretion by 25% ($P < 0.05$; $n = 6$). Dexamethasone in combination with insulin produced a 22% increase in leptin mRNA abundance relative to insulin ($P < 0.05$; $n = 4$), and T₃ stimulated a 28% increase in insulin-induced leptin mRNA abundance ($P < 0.05$; $n = 4$). Leptin receptor mRNA abundance was decreased by 25% with the combination of insulin and dexamethasone, relative to insulin-treated adipose tissue slices ($P < 0.05$; $n = 4$). Porcine GH decreased leptin receptor mRNA abundance by 17% ($P < 0.05$; $n = 6$). These data suggest that leptin secretion is a regulated phenomenon and that posttranslational processing may be significant. Alternatively, transport and exocytosis of leptin containing vesicles in the pig adipocyte may be quite complicated, which could account for the differences in observed mRNA abundance and protein secretion.

Key Words: Adipose, Leptin, Leptin Receptor, Swine

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Introduction

Leptin is a peptide secreted from the adipose tissue that may play an important role in suppressing appetite

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in swine through actions at the hypothalamus (Barb et al., 1998). Leptin mRNA has been detected in pig adipose tissue (Ramsay et al., 1998; Leininger et al., 2000), with expression exclusive to the adipocyte (Chen et al., 1997). Several studies have been published that demonstrate hormonal regulation of porcine leptin mRNA levels in vitro (Chen et al., 1997, 1998; Ramsay and White, 2000). Those studies have used adipocytes differentiated in vitro from neonatal adipose tissue, rather than adipocytes obtained from the pig. The current study was performed to examine the response of the leptin gene in porcine adipose tissue to hormonal stimuli.

Changes in leptin mRNA level were correlated with adiposity (Leininger et al., 2000). Differences in tissue leptin mRNA abundance were associated with differences in serum leptin concentrations (Ramsay et al., 1998). Secretion of leptin into the bloodstream was con-

firmed by western analysis (Ramsay et al., 1998) and by RIA (Barb et al., 2001). However, the relationship between changes in leptin mRNA abundance and leptin protein secretion from the adipose tissue in swine is unknown and was the purpose of the current study.

Materials and Methods

Yorkshire gilts (approximately 150 kg BW) were used for these experiments, with four to six gilts used per experiment. Gilts were provided water and a corn-soybean diet (18.3% CP, 3,059 kcal/kg of ME; as fed) ad libitum. Animals of this size were intentionally used because of the enlarged adipose mass containing a higher proportion of large adipocytes than present in the adipose tissue from smaller swine (Etherton et al., 1981). Adipose tissue with large adipocytes has been shown to express higher levels of leptin than tissue containing small adipocytes (Van Harmelen et al., 1998), thereby enhancing the ability to detect secreted leptin. Dorsal s.c. adipose tissue samples from between the second and fourth thoracic vertebrae were acquired following slaughter by electrical stunning and exsanguination according to procedures approved by the Institutional Area Animal Use and Care Committee. The middle s.c. adipose tissue was dissected free of the outer s.c. adipose to reduce variability in hormone responses. Dissected adipose tissue was diced into 1- × 4-cm strips and placed in Hanks buffer (37°C, pH 7.4) in screw-capped polypropylene Erlenmeyer flasks for transport to the laboratory, approximately 2 min from the abattoir.

In the laboratory, adipose tissue strips were placed in fresh Hank's buffer (37°C, pH 7.4). Adipose tissue strips were dissected clean of any extraneous muscle tissue and further separated into 1-cm cubes in a laminar-flow hood. Adipose tissue explants (approximately 100 mg) were prepared by slicing tissue cubes with a Stadie-Riggs microtome. Tissue slices (400 μ m thickness) were rinsed twice with fresh Hanks buffer (37°C, pH 7.4), blotted free of excess liquid, and weighed. Tissue slices were then transferred to 12-well tissue culture plates containing 1 mL of media 199 with 25 mM HEPES, 0.5% BSA, pH 7.4, and the various hormone supplements of interest. Triplicate tissue slices were incubated with either basal medium or hormone supplemented media in a tissue culture incubator at 37°C with 95% air:5% CO₂.

Hormones included 100 nM insulin, 100 ng/mL porcine GH, 1 μ M dexamethasone, 10 nM triiodothyronine, 250 ng/mL of IGI-1, or 100 ng/mL porcine leptin. Sterile hormone solutions were prepared and frozen in vials before initiation of the experiment. Porcine insulin (Sigma-Aldrich, St. Louis, MO) was solubilized in 0.001 N HCl. Dexamethasone (Sigma-Aldrich) was solubilized in ethanol. Porcine GH (USDA-pGH-B-1, Beltsville, MD) was diluted in 25 mM sodium bicarbonate buffer, pH 9.4 (Na₂CO₃/NaHCO₃). Triiodothyronine (T₃; Sigma-Aldrich) was prepared in 0.01 N NaOH. Human

recombinant IGF-I (Sigma-Aldrich) was solubilized with 0.1% acetic acid. Recombinant porcine leptin was acquired from A. Gertler (Raver et al., 2000) and solubilized in PBS. Following initial solubilization, all hormones were subsequently diluted in 0.5% BSA in saline, aliquoted, and frozen. Individual hormone aliquots were thawed for each day of use and diluted in incubation medium to the appropriate concentration. Control incubations in basal medium included a similar amount of vehicle to exclude the vehicle as a variable. The selected hormone concentrations were determined in preliminary experiments or based on concentrations described in the literature to affect leptin/leptin receptor expression for insulin (Ramsay and White, 2000), dexamethasone (Murakami et al., 1995), GH (Chen et al., 1998), IGF-I (Morash et al., 2000), T₃ (Yoshida et al., 1997), and leptin (Wang et al., 1999).

Following 24 h of incubation, tissue samples from these incubations were blotted and transferred to microfuge tubes with subsequent freezing in liquid N and storage at -80°C before analysis for mRNA abundance. Media from incubations was collected in microfuge vials and stored at -20°C before analysis for leptin content. Media samples were centrifuged (10,000 × g) for 10 min at 4°C and duplicate 100- μ L aliquots were collected for measurement of media leptin content by RIA (Multispecies RIA kit, Linco, St. Charles, MO). Human leptin standards were replaced with recombinant pig leptin standards. Recombinant pig leptin was obtained from A. Gertler (Raver et al., 2000). Cross-reactivity for the recombinant porcine leptin in the multispecies leptin RIA was 58%. The intraassay and interassay CV were 8.32 and 16.3%.

Gene Expression Analyses by Reverse-Transcription Polymerase Chain Reaction

Total RNA was isolated using TRI reagent according to the manufacturer's protocol (Sigma-Aldrich). Integrity of RNA was assessed via agarose gel electrophoresis and RNA concentration and purity were determined spectrophotometrically for absorbance of light at 260 and 280 nm. Reverse-transcription (RT) PCR (20 μ L) consisted of 1 μ g of total RNA, 50 U of SuperScript II reverse transcriptase (Invitrogen/Life Technologies, Carlsbad, CA), 40 U of an RNase inhibitor (Invitrogen/Life Technologies), 0.5 mmol/L dNTP, and 100 ng of random hexamer primers. Polymerase chain reactions were performed in 25 μ L containing 20 mmol/L Tris-HCl, pH 8.4, 50 mmol/L KCl, 1.0 μ L of the RT reaction, 1.0 U of Platinum Taq DNA polymerase (Hot Start; Invitrogen/Life Technologies), 0.2 mmol/L dNTP, 2.0 mmol/L Mg²⁺ (Invitrogen/Life Technologies), 10 pmol each of the leptin and leptin receptor specific primers, and 10 pmol of an appropriate mixture of primers and competitors specific for 18S rRNA (QuantumRNA Universal 18S Internal Standard; Ambion, Inc; Austin, TX). Thermal cycling parameters were as follows: 1 cycle 94°C for 2 min, followed by 30 cycles, 94°C for 30 s,

58°C for 30 s, and 72°C for 1 min, with a final extension at 72°C for 8 min.

The following primers were used for generating 348-bp PCR products corresponding to a portion of the pig leptin coding sequence: 5'-TGACACCAAAACCCTCATCA-3' (forward), 5'-GCCACCACCTCTGTGGAGTA-3' (reverse). The primers for the leptin receptor were used to generate a 393-bp product corresponding to the extracellular domain, thus encompassing the total leptin receptor population: 5'-ACTGGAGCACCCCCTTTACT-3' (forward), 5'-TGGTTGACCATCTGCAAGTC-3' (reverse).

The RT-PCR techniques were optimized in a series of preliminary experiments. The two-step duplex RT-PCR for leptin + 18S or leptin receptor + 18S were optimized for linearity (exponential amplification) from >25 to <35 cycles under the conditions described above.

Capillary Electrophoresis with Laser-Induced Fluorescence Detection

Aliquots (2 μ L) of RT-PCR samples were diluted 1:100 with deionized water before capillary electrophoresis with laser-induced fluorescence detection (CE/LIF). A detailed description and validation of the CE/LIF technique used in this study for quantitative analysis of RT-PCR products was reported previously (Richards and Poch, 2002). Briefly, a P/ACE MDQ CE instrument (Beckman Coulter, Fullerton, CA) equipped with an argon ion LIF detector was used. Capillaries were 75 μ m i.d. \times 32 cm μ SIL-DNA (Agilent Technologies, Folsom, CA). Enhance dye (Beckman Coulter) was added to the DNA separation buffer (Sigma-Aldrich) to a final concentration of 0.5 g/L. Samples were loaded by electrokinetic injection at 3.5 kV for 5 s and run in reverse polarity at 8.1 kV for 5 min. Integrated peak area for the PCR products separated by CE was calculated using P/ACE MDQ software (Beckman Coulter).

Quantification of Gene Expression

For this study, estimates of the abundance of mRNA were determined as the ratio of integrated peak area for each individual gene PCR product relative to that of a coamplified 18S internal standard (QuantumRNA Universal 18S Internal Standard; Ambion, Inc, Austin, TX). Standardization with a coamplified 18S internal standard will correct for sample handling. Thus, a corrected or relative value for the target gene sequence is produced for each sample. Although CE/LIF is a quantitative procedure for measuring RNA products (Richards and Poch, 2002), RT-PCR can best be described as semi-quantitative (Joyce, 2002; O'Connell, 2002). Relative values are presented as the mean \pm SEM of four to six individual determinations.

Statistical Analyses

The experimental model for these experiments was a completely randomized design. Data were normalized

relative to insulin medium (100 nM) to account for culture-to-culture variation. Data were analyzed by one-way AOV using SigmaStat software (SPSS Science, Chicago, IL). Mean separation was analyzed using Student-Newman-Keuls test. Means were defined as significantly different at $P < 0.05$.

Results

Leptin Secretion

Leptin secretion occurred during a 24-h exposure to the hormones and hormone combinations of interest. Basal secretion of leptin varied among the tissue cultures, ranging from 21 to 38 ng/mL for each 100 mg of tissue (29.7 ± 2.8 ; $n = 6$). Originally, the data were to be expressed relative to the basal medium; however, degradation of the mRNA was detected in this control group incubated with basal medium, indicating tissues incubated without a hormone supplement were deteriorating. Therefore, the data was expressed relative to the insulin treatment group wherein media leptin levels ranged from 26 to 39 ng·mL⁻¹·100 mg tissue⁻¹ (31.9 ± 3.2 ; $n = 6$), which did not differ significantly from the basal secretion rate (29.7 ± 2.8 ng·mL⁻¹·100 mg of tissue⁻¹; $P = 0.56$; $n = 6$). As a result of this variability between tissue cultures, data were expressed relative to leptin secretion in insulin treated cultures for each animal.

In the first experiment, dexamethasone (1 μ M) was tested in combination with insulin for its efficacy in stimulation of leptin secretion (Figure 1). The combination of insulin and dexamethasone stimulated leptin secretion into the medium by 60% ($P < 0.001$). In a second experiment, porcine growth hormone addition to insulin-containing medium inhibited leptin secretion by 25% ($P < 0.01$) relative to incubations containing insulin. Triiodothyronine (10 nM) had no effect in combination with insulin on leptin secretion from adipose tissue slices ($92.7 \pm 10\%$ of control; $P = 0.55$). Addition of IGF-I (250 ng/mL medium) in combination with insulin to the incubation medium did not affect leptin secretion ($83.7 \pm 11\%$ of control; $P = 0.39$).

Leptin Gene Expression

Following the experiments examining the regulation of leptin secretion, further experiments were performed to evaluate hormonal regulation of leptin gene expression using tissue samples from additional pigs. Leptin mRNA was normalized to 18S rRNA, and then data were expressed relative to tissue cultures incubated with insulin containing medium to account for the culture-to-culture variation.

The combination of insulin with dexamethasone produced a 22% increase in leptin mRNA abundance relative to insulin (Figure 2; $P < 0.01$; $n = 4$). Incubation of tissue cultures with porcine growth hormone and insulin for 24 h had no effect on leptin mRNA abundance

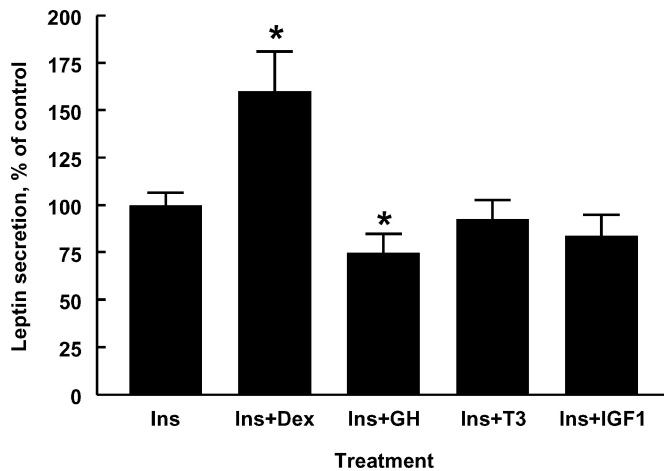


Figure 1. Relative percentage of leptin secretion in response to hormone exposure. Primary cultures of porcine adipose tissue were incubated with insulin (Ins, 100 nM), the combinations of insulin and dexamethasone (Dex, 1 μ M), insulin and porcine growth hormone (GH, 100 ng/mL), insulin and triiodothyronine (T3, 10 nM), or insulin and IGF-I (250 ng/mL) for 24 h, followed by collection of culture media for analysis of leptin content by RIA. Data are expressed relative to cultures incubated with insulin. Asterisks indicate that means differ ($P < 0.05$) from treatment with insulin ($n = 6$).

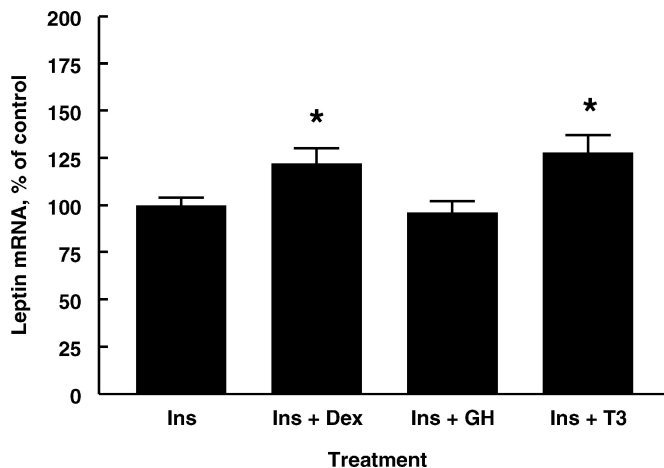


Figure 2. Relative abundance of leptin mRNA in response to hormone exposure. Primary cultures of porcine adipose tissue were incubated with insulin (Ins, 100 nM; $n = 6$), the combinations of insulin and dexamethasone (Dex, 1 μ M; $n = 4$), insulin and porcine growth hormone (GH, 100 ng/mL; $n = 6$), or insulin and triiodothyronine (T3, 10 nM; $n = 4$) for 24 h, followed by extraction for total RNA and subsequent reverse-transcription PCR analysis for abundance of leptin mRNA. Data are expressed relative to cultures incubated with insulin. Asterisks indicate that means differ ($P < 0.05$) from treatment with insulin.

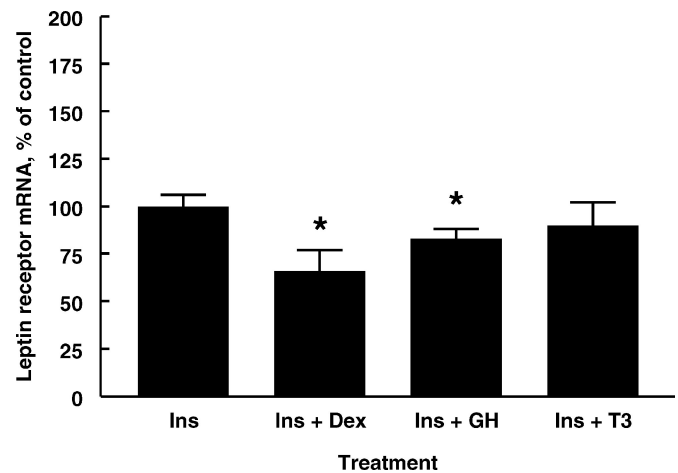


Figure 3. Relative abundance of leptin receptor mRNA in response to hormone exposure. Primary cultures of porcine adipose tissue were incubated with insulin (Ins, 100 nM; $n = 6$), the combinations of insulin and dexamethasone (Dex, 1 μ M; $n = 4$), insulin and porcine growth hormone (GH, 100 ng/mL; $n = 6$), or insulin and triiodothyronine (T3, 10 nM; $n = 4$) for 24 h, followed by extraction for total RNA and subsequent reverse-transcription PCR analysis for abundance of leptin receptor mRNA. Data are expressed relative to cultures incubated with insulin. Asterisks indicate that means differ ($P < 0.05$) from treatment with insulin.

($96.3 \pm 6\%$ of control; $P = 1.00$; $n = 6$), relative to cultures incubated with insulin alone. Tissue slices incubated with T₃ and insulin expressed leptin mRNA at a 28% higher level than slices incubated with insulin ($P < 0.01$; $n = 4$). Leptin in combination with insulin and dexamethasone produced a similar change in leptin mRNA level as insulin in combination with dexamethasone ($108 \pm 7\%$ of insulin + dexamethasone; $P = 0.38$; $n = 6$; data not presented).

Leptin Receptor mRNA Abundance

Following 24 h of incubation, insulin in combination with dexamethasone produced a 25% decrease in leptin receptor mRNA abundance (Figure 3; $P < 0.03$; $n = 4$). Growth hormone inhibited insulin induced leptin receptor mRNA abundance by 17% ($P < 0.001$; $n = 6$). The combination of insulin and T₃ produced a similar effect on leptin receptor mRNA abundance as insulin alone ($88.9 \pm 11\%$ of control; $P = 0.209$; $n = 4$). The addition of leptin in combination with insulin and dexamethasone had no effect on the abundance of leptin receptor mRNA ($99.3 \pm 9\%$ of insulin + dexamethasone; $P = 0.710$; $n = 6$; data not presented).

Discussion

Adipose tissue is the primary source of leptin secreted into the bloodstream (Harris, 2000). This secretion is

through both regulated constitutive secretory pathways involving intracellular vesicle transport (Bradley et al., 2001). Hormonal regulation of leptin secretion has been demonstrated for both human and rodent adipose tissue (Guerre-Millo, 2002). Although regulation of leptin gene expression has been evaluated in newly differentiated pig adipocyte cell cultures (Chen et al., 1998), regulation of the secretion of porcine leptin *in vitro* has not been previously examined.

Insulin has been reported to stimulate leptin secretion from human and rodent adipocytes at concentrations as low as 0.16 nM (Mueller et al., 1998; Russell et al., 1998). The current study could not replicate these results with a concentration of 100 nM insulin. This may be the consequence of species variation due to the relative insulin resistance of the pig adipocyte compared with human or rodent adipocyte (Mersmann, 1989).

Because the insulin treatment group was used to make all comparisons of mRNA abundance, we cannot discern from the results of this study whether insulin stimulates or inhibits leptin gene expression, as it served as the control. However, previous research using pig adipocytes formed *in vitro* (Chen et al., 1998; Ramsay and White, 2000) and human adipose tissue (Russell et al., 1998) demonstrated that insulin can stimulate leptin expression, although the results were not usually evident at 24 h of incubation, but only after longer periods of time (e.g., 72 to 96 h).

Glucocorticoids have been previously demonstrated to induce leptin mRNA abundance and protein secretion from human (Halleux et al., 1998; Russell et al., 1998) and rodent adipose tissues (Murakami et al., 1995; Sliker et al., 1996). The increase in leptin mRNA steady-state level may be due in part to enhanced transcription, probably through binding of the steroid-receptor complex to the glucocorticoid response element in the regulatory region of the leptin gene (Gong et al., 1996). In the current study, the combination of insulin and dexamethasone stimulated leptin gene expression and protein secretion. Kanu et al. (2003) reported that insulin is required for a dexamethasone stimulation of leptin secretion from isolated subcutaneous adipocytes, whereas Halleux et al. (1998) indicated that insulin decreases dexamethasone-induced increases in leptin mRNA abundance and protein secretion from omental adipocytes. These conflicting results may be the consequence of regional differences in adipose tissue or methodology and demonstrate the need for examination of leptin regulation in the different adipose tissues of the pig.

The majority of research on the porcine leptin receptor has focused primarily on detection of the receptor rather than regulation of the receptor (Czaja et al., 2002). Lin et al. (2003) have recently reported that GHRH inhibited leptin receptor expression in the porcine pituitary *in vitro*. Unfortunately, few studies on the regulation of leptin receptor gene expression in the adipocyte have been performed. The data from the cur-

rent study indicate that dexamethasone can inhibit total leptin receptor gene expression. Smith and Waddell (2002) have previously reported that dexamethasone can inhibit leptin receptor expression in the placenta, although no mechanism was defined. The inhibition of leptin receptor expression by dexamethasone, while stimulating leptin gene expression and protein secretion, might imply a downregulation of the receptor in the presence of elevated leptin protein levels, although this is speculation without examination of the leptin receptor binding and turnover.

Porcine GH inhibited insulin-induced leptin secretion, indicating some species variation in the regulation of leptin secretion. Growth hormone promotes insulin resistance in swine adipose tissue (Walton and Etherton, 1986). This may play a significant role in the observed inhibition of insulin induced leptin secretion because GH seems to not have a direct effect on the secretion of leptin as suggested by studies with mouse adipocytes (Fain et al., 1999; Lee et al., 2001). The inhibition of leptin secretion in the current study agrees with experiments in humans demonstrating an inverse relationship between plasma GH and plasma leptin concentrations (Florkowski et al., 1996; Fors et al., 1999). Although these *in vivo* experiments did not measure insulin, they cannot exclude the possible interaction of growth hormone with insulin mediated leptin secretion.

In vivo GH administration has been demonstrated to inhibit leptin gene expression by 56% in swine adipose tissue (Spurlock et al., 1998). *In vitro*, GH can inhibit the insulin-induced expression of leptin in bovine adipose tissue (Houseknecht et al., 2000) or porcine adipocytes differentiated in culture (Chen et al., 1998). In the current study, we did not observe an attenuation of the response to insulin when GH was added to the medium. The various responses among the reports suggest that species-specific or age-related responses to GH by the leptin gene exist. The data from the leptin secretion experiment in the current study indicate that GH inhibits leptin secretion, yet no change in leptin mRNA level was detected. These data would imply that posttranslational processing of leptin may be significant in the pig adipocyte. Alternatively, transport and exocytosis of leptin containing vesicles in the pig adipocyte may be quite complicated. Little is known concerning the mechanisms regulating the transport and exocytosis of these vesicles (Bradley et al., 2001). The disagreement between the leptin mRNA and protein secretion response to GH demonstrates the risk of interpreting mRNA data as indicative of an endocrine response to a stimuli, when it is only indicative of the mRNA.

Growth hormone was able to inhibit the insulin-induced expression of the leptin receptor. No specific effect of GH on the leptin receptor has been reported. This inhibitory effect may be the indirect through GH antagonism of insulin action in swine adipose tissue, as previously reported (Walton and Etherton, 1986).

The relative role of changes in the level of leptin receptor mRNA to either number or activity of functional leptin receptors has not been elucidated. There is a critical need for examination of leptin receptor kinetics as the potential significance of leptin receptor mRNA data has not been confirmed.

Many of the effects of GH have now been attributed to IGF-I through paracrine induction. Porcine GH has been demonstrated to produce a rapid increase in IGF-I expression in swine adipose tissue (Ramsay et al., 1995). In the current study, IGF-I did not alter leptin secretion into the medium, unlike GH, which inhibited leptin secretion by 25%. Isozaki et al. (1999) reported lower serum leptin concentrations and higher IGF-I levels in acromegalics than in normal subjects, suggesting that GH directly interacts with adipose tissue to decrease leptin expression. However, a significant positive correlation between serum leptin and serum IGF-I has been reported for normal human subjects, but was not associated with IGF-I regulation of leptin synthesis or secretion (Gomez et al., 2003). These studies support the hypothesis that GH is acting directly on the adipocyte and not through IGF-I to regulate leptin secretion.

Fain and Bahouth (1998) reported that T_3 treatment of adipose tissue explants from hypothyroid rats produces an increase in leptin secretion and elevated leptin mRNA levels when used in combination with insulin and dexamethasone. In the current study, T_3 had no effect on leptin secretion from porcine adipose tissue explants in the presence of insulin, but T_3 increased leptin mRNA abundance by 28%. Treatment with T_3 alone has been shown to increase leptin mRNA and secreted leptin levels in cultures of 3T3-L1 adipocytes (Yoshida et al., 1997), to have no effect in human omental adipose tissue explants (Menendez et al., 2001), or to inhibit leptin secretion and gene expression in human s.c. adipose tissue (Kristensen et al., 1999). The true function of T_3 in regulation of leptin gene expression and secretion may be difficult to determine, as the variability between studies suggests that the response depends on the entire hormonal milieu, time of incubation, tissue source, and species. Alternatively, variable conditions for culture among the studies might contribute to the differences in the T_3 response.

Implications

Leptin is a hormone produced by pig adipose tissue that can affect feeding behavior, animal health, and reproduction. Studies in pigs have demonstrated that leptin can decrease feed intake. This experiment was designed to determine whether the production and release of porcine leptin is a regulated phenomenon. Secondly, this study attempted to determine whether expression of leptin and its receptor could be manipulated by hormonal mechanisms in mature animals. The data demonstrate that leptin secretion can be both stimulated and inhibited. Secondly, expression of the leptin

gene and the leptin receptor gene are responsive in adult swine. These results suggest the potential to manipulate the efficiency of feeding behavior in finishing animals at a time when changes in body composition can be critical for improving profitability.

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